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The morphology, physiology, and fine structure of a toluene-oxidizing strain of *Pseudomonas putida*

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AN ABSTRACT OF THE THESIS OF Barry Clayton Anderson for the
Master of Science in Biology presented September 18, 1992.

Title: The Morphology, Physiology, and Fine Structure of a
Toluene-Oxidizing Strain of *Pseudomonas putida*.

APPROVED BY THE MEMBERS OF THE THESIS COMMITTEE:


Mary L. Taylor, Chair


Lester J. Newman


David T. Clark


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Gordon L. Kilgour

The role of microorganisms in the degradation of
xenobiotics in the environment is well established. Bacteria
from the genus *Pseudomonas* are particularly well adapted to
the degradation of hydrocarbons, aromatics, and numerous
other natural and introduced substrates. We have isolated a

strain of *Pseudomonas putida*, designated PC2P15, that uses toluene, phenol, benzene, and a number of other substrates as its sole sources of carbon and energy.

Morphological, biochemical, and metabolic analysis of PC2P15 has revealed significant changes in the fatty acid content, the presence of extracellular membrane vesicles, the possible production of an emulsifying agent and the apparent simultaneous use of multiple metabolites in bacteria exposed to toluene, succinate, or toluene and succinate when compared to those organisms growing in sucrose alone. ^{14}C -sucrose uptake in bacteria exposed simultaneously to both sucrose and toluene or sucrose and succinate each showed an approximately 50% decrease in ^{14}C -sucrose uptake when compared to organisms growing in sucrose alone. Organisms exposed to a combination of sucrose, toluene, and succinate showed an approximately 75% drop in ^{14}C -sucrose uptake when compared to organisms grown in sucrose alone. The addition of cyclic-AMP had no effect on ^{14}C -sucrose uptake for any combination of substrates.

THE MORPHOLOGY, PHYSIOLOGY, AND FINE STRUCTURE
OF A TOLUENE-OXIDIZING STRAIN
OF *PSEUDOMONAS PUTIDA*

by

BARRY CLAYTON ANDERSON

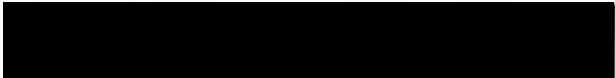
A thesis submitted in partial fulfillment of
the requirements for the degree of

MASTER OF SCIENCE
in
BIOLOGY

Portland State University
1992

TO THE OFFICE OF GRADUATE STUDIES:

The members of the committee approve the thesis of
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DEDICATION

I dedicate my research and this Thesis to my wife, Claire Roff Anderson, whose love and unwavering support have allowed me to pursue my dreams.

I also dedicate this Thesis to my parents, Virginia Hyatt Anderson and Wallace Julian Anderson, for their love and continued belief in me throughout my life and in the midst of many storms.

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I would like to acknowledge Dr. Mary L. Taylor of the Portland State University Department of Biology for her help and encouragement in the preparation of this Thesis and for her intellectual contributions to the research upon which this Thesis is based.

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INTRODUCTION

BIOREMEDIATION

Increasing contamination of soils and water with toxic pollutants including the seepage of oil (Figure 1) has greatly increased interest in the role of microbial communities in the degradation and bioremediation of toxic organic molecules and various recalcitrant aromatic compounds. This problem was brought to the forefront on 24 March 1989 when the oil tanker Exxon Valdez ran aground in Prince William Sound, Alaska. Massive cleanup efforts included the use of bacteria to degrade a portion of the eleven million gallons of crude oil spilled into the previously pristine Sound. Initially, some success was achieved through the use of both water soluble and oleophilic fertilizers to enhance bioremediation using naturally occurring soil organisms (1,40,60). Despite these promising initial results, many of the factors governing *in situ* bioremediation remain unknown.

A number of parameters must be considered when inoculating a culture enriched for a particular hydrocarbon into a contaminated site. The success of the bioremediation effort may be influenced by predation and competition or by abiotic stresses for introduced organisms such as low

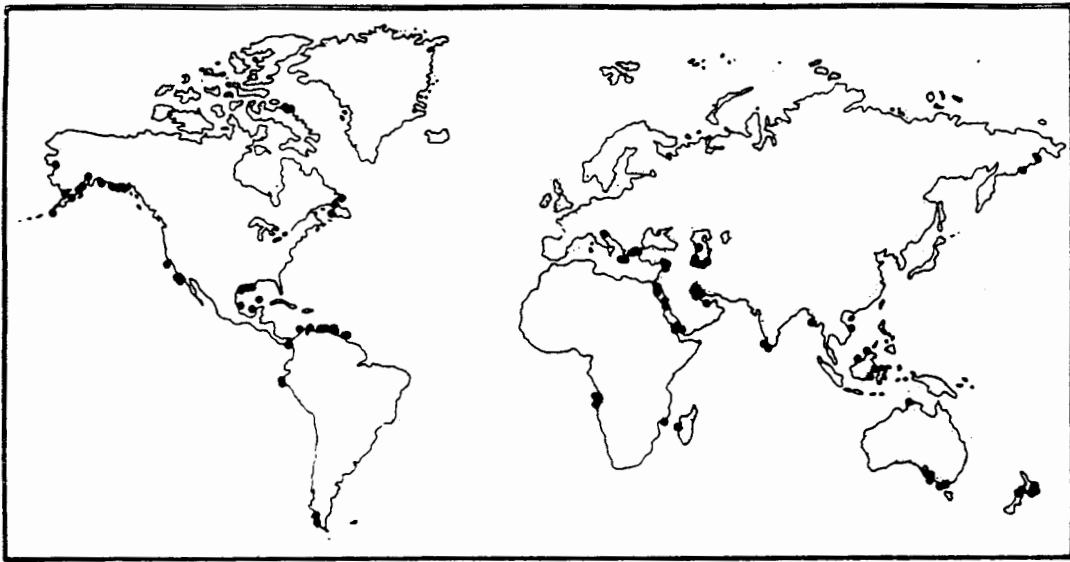


Figure 1. Locations of marine oil seeps. (*Oil in the Sea: Inputs, Fates, and Effects*, 1985. p.47 (63).)

nutrient levels, high contaminant levels, toxins, or the preferential use of other organic compounds (57).

Many investigators are attempting to understand the basic physiology and metabolism of those microorganisms capable of degrading hydrocarbons both *in situ* and in the laboratory (10,11,12). A recent study by Hill (2) indicated the presence of *Alcaligenes* A5 and a number of other biodegradative organisms in an area contaminated with 4-chlorobiphenyl. Most probable number for growth on polychlorinated biphenyl (PCB) followed by gene probe analysis revealed the presence of the 4CB plasmid, which codes for the degradation of PCB, scattered throughout the indigenous populations.

In another study, enrichment isolates of selected strains of *Pseudomonas* have been used successfully for *in situ* bioremediation of hydrocarbon-contaminated soil and groundwater in California (3). Some investigators have shown that, in sites where long standing contamination has occurred, populations of soil organisms have adapted to degrade alkanes and aromatics. A recent site survey revealed the presence of 244 gasoline-degrading isolates in an aquifer contaminated with unleaded gasoline (4). The four major genera represented at this site were *Pseudomonas*, *Alcaligenes*, *Micrococcus*, and *Nocardia*. Madsen (41) has reported the biodegradation of naphthalene *in situ* in a contaminated aquifer. A recent report indicated that a

luciferase marker has been placed in a strain of *P. fluorescens* and currently is being tested as a bioluminescent indicator for naphthalene exposure and degradation (42). Recent reports (58) have shown that there are unanticipated interactions of substrates, products, and organisms in a benzene, toluene, and xylene (BTX) contaminated aquifer which affected both the indigenous and introduced populations. This finding reinforces the fact that bioremediation in natural systems is a complex, multifaceted process in which indigenous or introduced organisms react in ways that are not always predictable from studies done in the laboratory.

Other investigators have reported that a wide variety of compounds are being degraded by naturally occurring soil and water organisms in sites of long standing contamination. Among the compounds that have been found to be degraded are benzoate (44), *p*-nitrophenol (45), chlorinated biphenyl (46), 1,2,3,-trichloro- and 1,2,4,5,-tetrachlorobenzene (59), and others.

The rapidly growing field of recombinant DNA technology is being exploited in the biodegradation of xenobiotics and toxic waste and shows some promise, although this type of research applied *in situ* is closely monitored by the EPA (43). Some research has shown, however, that organisms inoculated into contaminated sites are not able to pass on the plasmids containing the genes for the metabolism of

hydrocarbons with any degree of efficiency in soils. As one might expect, intrageneric transfer is the most efficient, with approximately 10^{-2} transfers per transconjugate. However, intergeneric plasmid transfers are reported to be as low as 10^{-7} transfers per transconjugate (47). The authors contend that these low figures of 1 plasmid transfer per 100 intragenerically and 1 transfer per 10,000,000 intergenerically may not be abnormal *in situ* due to the interplay of a multitude of abiotic factors in soils and water.

It has been determined (5) that the rate of degradation of hydrocarbons and organic solvents *in situ* is controlled by a number of complex factors including concentration of the contaminant(s), temperature, available nutrient levels, emulsification, salinity, and the condition of the soil or water at the site.

Currently, the economic feasibility of microbially mediated bioremediation, microbially enhanced oil recovery, and even the use of bacteria in coal bioprocessing is an area of intense research (6,7). Among the bacteria being investigated, those belonging to family *Pseudomonadaceae* are some of the most promising because of their impressive metabolic capabilities.

THE PSEUDOMONADS

den Dooren de Jong in 1926 listed 80 organic compounds used by a species of *Pseudomonas* (as part of the research for his doctoral dissertation) (10). Other investigators have extended the scope of our knowledge of the physiology and metabolic capabilities of the pseudomonads (20,21). Most of the members of this genus grow quite rapidly on a variety of metabolites, exhibiting growth typical of r-type selection. In the natural environment, r-type organisms with the nutritional versatility of the pseudomonads would be expected to respond to the sudden influx of a new carbon source by outcompeting the more exacting K-type organisms. It is this very versatility that makes the pseudomonads excellent subjects for research not only into their role in bioremediation, but also into the biochemistry, physiology, and genetics of this and other groups of microorganisms in the environment.

Organisms in genus *Pseudomonas* are gram negative, motile rods. Typically, they display one or more polar flagella. They are absolute respirers and are usually oxidase positive. Pseudomonads never ferment. Some of the members of this group synthesize a yellow-green water soluble pigment that characteristically fluoresces under ultra-violet light. Members of this genus are common, free-living organisms in soils, fresh or marine waters, and in

association with plants and animals. With a few exceptions, such as certain strains of *P. aeruginosa*, most members of this genus are not pathogenic. Pseudomonads are capable of great metabolic diversity and are able to use a large variety of organic compounds as sources of carbon and energy. Some strains may utilize as many as one hundred different compounds including sugars, amino acids, hydrocarbons, aromatics, and fatty acids.

BIOCHEMISTRY OF AROMATIC DEGRADATION

Most naturally formed mono- and polycyclic aromatic compounds result from forest fires, from the combustion of fossil fuels, and from the degradation of lignin by fungi (8,9). It is not surprising that many microorganisms have adapted metabolic pathways capable of degrading these substances, especially when one considers the total amount of hydrocarbons and other organic substances being introduced into the environment every year. For example, it has been estimated that more than one billion gallons of toluene are produced each year by the industrialized nations. This production is accompanied by the spills, dumping, and accidents that are the seemingly inevitable results of human activity.

The benzene nucleus is exceedingly abundant in the biosphere and, because of its resonance structure, is

extremely stable. Most of the degradation of the benzene nucleus is done by microorganisms (14,15).

As shown in Figure 2, the prerequisite reaction for the breaking of the benzene nucleus is either a single or double hydroxylation of the ring depending on the substituents present. It was originally believed that the product of the double hydroxylation of benzene was a trans-benzene dihydrodiol (22,23). However, further investigation (16) revealed that a strain of *Pseudomonas* growing rapidly on toluene would oxidize benzene and catechol, but that trans-benzene dihydrodiol was not oxidized at a substantial rate. Other investigators (17,18) showed that a cis-benzene dihydrodiol was formed prior to the formation of catechol and that molecular oxygen was necessary for the hydroxylation reaction. Final proof of cis-benzene dihydrodiol as the pre-catechol intermediate was provided by researchers (18,39) using a mutant strain of *P.putida* deficient in cis-benzene dihydrodiol dehydrogenase.

There are various enzyme systems used by the pseudo-monads for the initial hydroxylation(s) of the benzene nucleus. Depending on the substituent groups present on the ring, the organisms may use either a monooxygenase system, a dioxygenase system, or both (33,35).

The dioxygenase system is controlled by the TOD plasmid. The dioxygenase is a three component enzyme system consisting of a flavoprotein reductase, a (2Fe-2S)

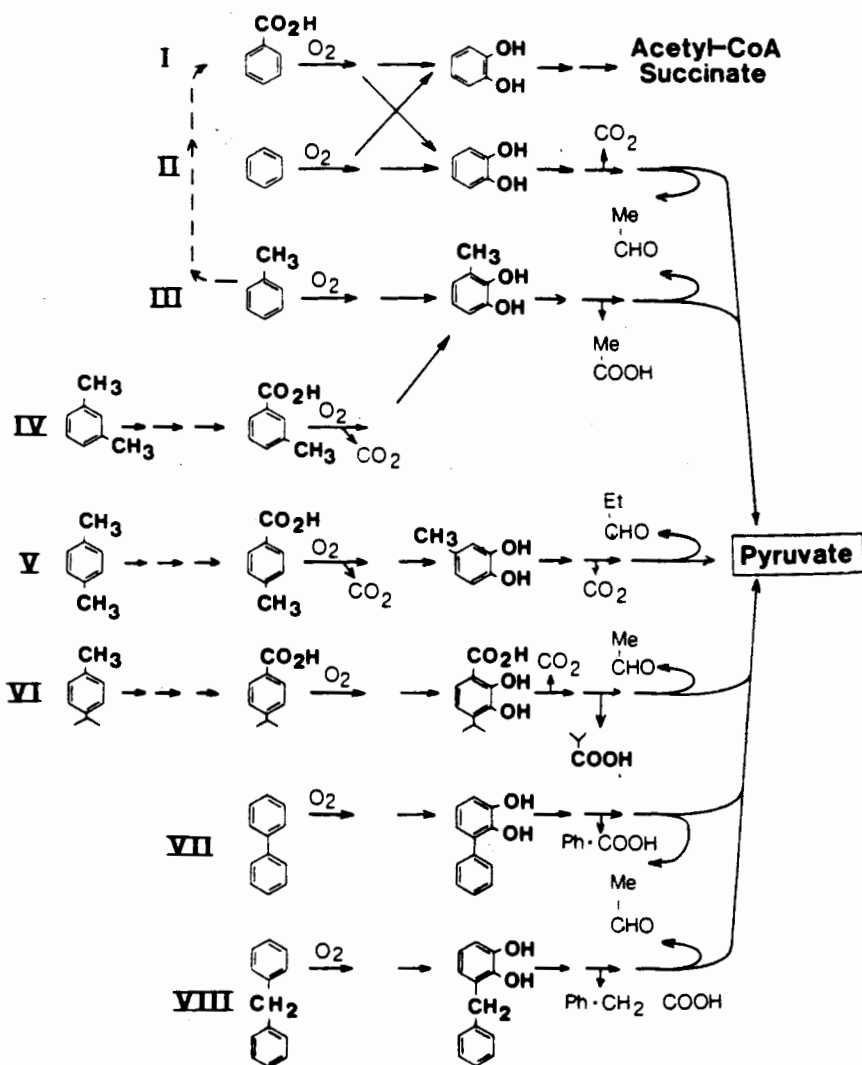


Figure 2. Summary of aromatic catabolic pathways. (From Dagley, S., 1986 (19).)

ferredoxin designatedferredoxin_{TOL}, and an iron-sulfur protein designated ISP_{TOL}. Electrons for this initial reaction, provided by NADH, are transferred to the flavoprotein and then subsequently passed to theferredoxin_{TOL}. The electrons are finally transferred to ISP_{TOL} and then to the cis-toluene dihydrodiol. The dioxygenase pathway forming cis-toluene dihydrodiol is followed by a dehydrogenation of the dihydrodiol by cis-toluene dihydrodiol dehydrogenase (36,37,38).

The monooxygenase is also a three component system (36) controlled by the TOL plasmid. The monooxygenase can form an o- or p-creosol, benzyl alcohol, or other intermediate as the initial product (36).

There are at least two major ways in which the benzene nucleus is cleaved following the formation of the catechol, protocatechuate, or other intermediate. Ortho-fission breaks the ring between the pair of hydroxyl groups which are almost always on adjacent carbons. Meta-fission opens the ring between an hydroxyl group and a non-hydroxyl substituent on the ring.

Ortho-fission is somewhat restricted in the degradation of hydrocarbons because of the seeming intolerance of the ortho-fission enzymes to the presence of substituent groups on the ring. Meta-fission enzymes generally tolerate the presence of methyl groups and other substituents, as is the case with the degradation of toluene.

In the case of benzene, ortho-fission of the catechol gives rise to cis, cis-muconic acid and then to beta-ketoadipate. Further reactions form the Krebs cycle intermediates succinate and acetyl-CoA. Meta-fission of the catechol leads to the formation of 2-hydroxy-cis, cis-muconic semialdehyde and then to acetoacetate, acetaldehyde, and pyruvate, and enters the Krebs cycle as acetyl-CoA (24).

RESEARCH FOCUS

Our laboratory has isolated a strain of *Pseudomonas putida* from the soil surrounding a bioremediation pond in Eugene, Oregon. This strain metabolizes a variety of straight and branched chain hydrocarbons and aromatics, including up to 50% (v/v) toluene as the sole sources of carbon and energy. Other investigators (13) have shown that certain strains of *P.putida* can withstand a 30-50% (v/v) toluene exposure and apparently thrive.

The focus of our research has been the investigation of how these organisms not only survive direct exposure to highly toxic organic solvents such as toluene, but how they are able to emulsify or partition the toluene molecule for use in their metabolism. We have asked how these solvents affect the outer membrane and have sought to discover the role of toluene exposure and metabolism in the physiological processes and fine structures of these bacteria. We have investigated changes in membrane fatty acid content, the

production of an emulsifying agent or agents, the significant depression of sucrose uptake in the presence of toluene and/or succinate, and the apparent simultaneous use of multiple metabolic pathways. We have looked at the relatively dramatic changes in their overall morphology and growth rate when compared to control organisms. We have also investigated the role of toluene as it relates to metabolic stress and compared our findings with the findings of other researchers studying the role of nutrient deprivation and metabolic stress on the physiology of microorganisms in the environment (30,31,32).

MATERIALS AND METHODS

ISOLATION, IDENTIFICATION, AND TESTING

Soil samples were taken from a bioremediation pond in Eugene, Oregon which was contaminated with various hydrocarbons including trichloroethylene. We used standard enrichment techniques to select an organism capable of growth in a modified Bushnell-Hass Medium with toluene as the sole source of carbon and energy.

The isolate was grown in 20ml of a slightly modified Bushnell-Hass (mBH) minimal salts media (pH 7) (Table I) with various hydrocarbons as the sole source of carbon and energy. All chemicals and reagents used were analytical or molecular biology grade unless otherwise indicated. Toluene was supplied to the organisms as the vapor in 125ml side-arm flasks. The bacteria were grown at room temperature and aerated by shaking at 100rpm on a New Brunswick Model S-3 rotating shaker. In tests requiring direct exposure to toluene as the liquid, the organisms were grown in 125ml Erlenmeyer flasks at room temperature and aerated at 100rpm. The exponentially growing cultures were passaged every 72 hours following a characteristic 48 hour lag phase.

Standard tests were performed to identify the isolate (Table II). These tests included a test for motility in

TABLE I
MODIFIED BUSHNELL-HASS FORMULATION

MgSO ₄ . 7H ₂ O.....	0.41g/L
CaCl ₂ . 2H ₂ O.....	0.04g/L
KH ₂ PO ₄	1.00g/L
K ₂ HPO ₄	1.00g/L
NH ₄ NO ₃	1.00g/L
FeSO ₄ *.....	0.05g/L
Resazurin#.....	0.002g/L
Phenol Red^.....	0.001g/L

* Change from Difco Formulation of FeCl₂
 # Optional Indicator of O₂
 ^ Optional Indicator of pH
 (made in sterile dd H₂O)

TABLE II
STANDARD TESTS TO IDENTIFY ISOLATE

Gram Stain.....	negative
Motility.....	positive
Oxidase.....	positive
Catalase.....	positive
O/F.....	respirer
Morphology.....	rod
Flagella.....	polar
Gelatinase.....	negative
Starch Hydrolysis.....	negative
Siderophore Production.....	positive
PHB Production.....	negative
Fluorescence.....	positive
Nitrate Reduction.....	negative
Nitrite Reduction.....	negative

GROWTH TEMPERATURES (in degrees centigrade)

Growth at 4 ⁰	negative
Growth at 10 ⁰	negative
Growth at 15 ⁰	slow positive
Growth at 20 ⁰	positive
Growth at 25 ⁰	positive
Growth at 32 ⁰	slow positive
Growth at 42 ⁰	negative

OTHER

Fatty Acid Analysis..... 81% positive
Pseudomonas putida

tubes of motility agar, gross colony morphology on plates of nutrient agar, Luria agar, TGYE agar, CAS agar (55) for the production of siderophores, and King's B agar for the test for fluorescent pseudomonads. The isolate exposed to toluene while growing on agar was grown on mBH with 1.5% DIFCO certified agar. The bacteria were streaked onto a sterile Nucleopore 0.22 μ m filter and placed in an enclosed glass tank with toluene vapors and incubated at room temperature. It should be noted that, in all cases when toluene was used as the substrate, the use of glass plates and other utensils was necessary due to the destructive effect of toluene on most plastics. The Nucleopore filter was not significantly affected by toluene during the required growth period. The isolate was tested with various hydrocarbons as the sole source of carbon and energy using the mBH minimal salts medium (Table III). Control organisms were grown in mBH with sucrose as the sole source of carbon and energy unless otherwise indicated.

GROWTH AND PROTEIN ANALYSIS

Growth rate studies were performed in 20ml mBH medium with toluene as the sole source of carbon and energy, incubated at room temperature and aerated at 100rpm as described above. Controls were grown in mBH medium with sucrose. Turbidity was measured on a Klett-Summerson Photoelectric Colorimeter using both a red and purple

TABLE III
SUBSTRATE UTILIZATION BY PC2P15

<u>Sugars and Other Metabolites</u>	<u>Lag Time*</u>	<u>Growth</u>
Glucose	3	+
Sucrose	<1	++++
Maltose	<1	+++
Galactose	<1	++++
Tryptose	<1	++++
Urea	<1	++++
AC Media	-	-
AC Media + Glucose	3	+
<u>Agars</u>		
TGYE	1	++++
Luria	1	+++
Nutrient	1	+++
Blood	1	+
<i>Pseudomonas</i> Isolation	1	++++
Soil Extract	1	++
Nutrient + 5% NaCl	-	-
<u>Hydrocarbons and Aromatics</u>		
Toluene	2	++++
p- and o-Xylene	3	++++
2,4-D	17	+++
2,4,5-T	-	-
Motor oil	6	+++
Diesel	6	+++
Unleaded gasoline	5	+++
Hexane	-	-
Dodecane	5	+++
Phenol 7% (v/v)	-	-
Phenol 2.5% (v/v)	12	+
Benzene	2	++++
Catechol 5% (v/v)	1	++++
Benzoic acid	8	++
Crude Oil	11	+++
1,3-DCE	-	-
p-amino salicylic acid	31	++++

*=approximate lag time in days:toluene inoculum
(as measured by turbidity or the appearance of colonies)

filter. Measurements were taken every two hours for the first eight hours, every four hours for the next sixteen hours, and every twelve hours thereafter for a total of nineteen readings. The growth rate study was performed twice and the figures averaged.

Fatty acid analysis and tentative identification of this isolate was kindly performed by Warren L. Landry of the Food and Drug Administration Laboratory in Dallas, Texas. The analysis used a Hewlett-Packard MIS GC and MIS 3.2 analytical software.

To obtain protein for the cytoplasmic protein analysis, exponentially growing organisms (approximately 10^7 cells/ml) were harvested at 72 hours post passage and centrifuged at 10,000xG for 30 minutes to pellet the cells. Cells were washed once in sterile 0.01M phosphate buffer (pH 7) and centrifuged at 10,000xG for ten minutes. The cells were resuspended in 2.5ml sterile 1mM HEPES (pH 7) and 2.5ml sterile 0.9% NaCl and placed on ice. The cells were broken in a chilled (4° C) Amicon French Pressure Cell at 28,000psi. The cell lysate was centrifuged at 3000xG for five minutes to remove unbroken cells. The supernatant was collected and concentrated at 4° C in Centriprep Protein Concentrators (Amicon) with a molecular weight cutoff of 10,000 daltons. This reduced the total supernatant volume from 5ml to 500 μ l. All concentration protocols were carried

out according to the manufacturer's directions. Concentrated cell lysate was placed on ice.

A 250ul aliquot of the sample was placed into 1.5ml microfuge tube to which 500ul of sample buffer was added. The sample buffer consisted of 4% sodium dodececyl sulfate (SDS), 12% glycerol, 2% 2-mercaptoethanol, 0.01% bromophenyl blue, 1mM phenylmethylsulfonylfluoride (PMSF) (Sigma), and 100mM dithiothreitol (DTT) (Sigma) in 50mM TRIS-HCl (pH 6.8). To reduce viscosity, 1mg/ml (final concentration) DNase I (Sigma) was added. The sample was incubated at room temperature for 30 minutes. After incubation, the sample was boiled for 3 minutes at 100° C and centrifuged for 2 minutes at 12,000xG. The sample was loaded onto a 12% discontinuous SDS polyacrylamide gel and electrophoresed at 260 volts for approximately three hours. The gel was fixed in 10% trichloroacetic acid (TCA) for 1 hour and stained with Coomassie R-250 for 24 hours.

Total cell membranes were prepared as previously described (25,26,27) and analyzed using the SDS polyacrylamide gel electrophoresis technique above with the exception that Triton X-100 was added to the membrane sample buffer to give a final concentration of 1%.

MICROSCOPIC ANALYSIS

Transmission electron microscopy was performed by Ken Tiekotter at the Good Samaritan Hospital Institute of Neuroscience, EM Research Laboratory, Portland, Oregon using a Zeiss EM10-CA transmission electron microscope. Light micrographs were obtained using a Zeiss compound microscope mounted with a Zeiss C-35 camera. Unless otherwise indicated, all light micrographs are at a magnification of 1200X.

METABOLIC ANALYSIS

Sucrose uptake studies were performed using the standard growth conditions described above. Each flask contained 20ml of mBH to which was added 2uCi of ^{14}C -sucrose along with 0.1% (final concentration) sterile unlabeled sucrose. To selected flasks, succinate (pH 6.8) was added at a final concentration of 0.1%. Cyclic-AMP was added to selected flasks at final a concentration of 1mM. The organisms were allowed to grow for 144 hours (unless otherwise specified) at room temperature with aeration at 100rpm. Harvested cells were washed twice in sterile 0.01M phosphate buffer (pH 7) before lysis in the French Pressure Cell as described above. One hundred microliters of bacterial cell extract was added to 10ml of scintillation fluid and counted using a Beckman Model LS-9000 liquid

scintillation counter. These experiments were performed six times and the data was averaged.

Colorimetric protein assays were performed using a Pierce BCA (Bicinchoninic Acid) Protein Assay Kit following the methods of Smith (28). In each assay, a 100ul aliquot of bacterial extract was diluted 1:10, placed in assay buffer per manufacturer's directions and read on an Ultraspec III Spectrophotometer (Pharmacia) at a wavelength of 562nm. Bovine Serum Albumin (BSA) was used as the standard.

Tests for the production of an emulsifying agent or surfactant was done according to the method of Jain (29). In this test, drops of suspended cells or pelleted cells of organisms grown in toluene or sucrose were placed on a sucrose, oil, or solvent-coated surface and observed for the collapse of the pellet or drop of suspended cells, indicating the production of an emulsifying agent or agents or the production of a surfactant.

RESULTS

IDENTIFICATION OF PC2P15

The soil sample from the contaminated bioremediation pond yielded 42 colony forming units (CFU) by standard plate count techniques. Each CFU was serially diluted in sterile water to approximately 10^{-8} and 1ml was inoculated into 10ml capped tubes with 9ml mBH. Each tube contained a different carbon source as shown in Table III. One isolate, (PC2P15), was found to utilize 13 of the 17 substrates tested.

Growth of *P.putida* PC2P15 on the various substrates was extremely variable, with lag times ranging from 1/2 hour with several sugars to 31 days with *p*-amino salicylic acid. (Table III). Concentration of the hydrocarbon was also found to be important. For those organisms from the isolated strain (PC2P15) growing on vapor at the liquid/vapor interface, concentration of the substrate was of no importance. For those organisms growing with the hydrocarbon added directly to the medium, high concentrations of some of the substrates was found to be inhibitory to growth. For example, toluene concentrations in excess of 50% (v/v) and phenol concentrations in excess of 5.0% (v/v) were found to totally inhibit growth. Growth on alkanes such as unleaded gasoline was not inhibitory up to a 50% (v/v) concentration.

Non-volatile substrates such as catechol or 2,4-D showed no differences in growth regardless of substrate concentration. Concentrations of up to 70% (v/v) were tried for all volatile and non-volatile hydrocarbons.

Following the substrate testing and limitations experiments, PC2P15 was grown primarily on toluene. Every 10 days the isolate was transferred to a substrate on which it had previously grown, but which required enzymes other than those in the toluene degradation pathway to be utilized. It was found that, in many cases, the organisms had either lost the ability to degrade the newly introduced substrate or needed an exceedingly long lag period before growth began.

GROWTH

Growth curves (Figure 3) are the mean of two identical experiments. Even with an inoculum of bacteria continuously grown in toluene, it can be seen that there was a lag time of approximately 48 hours for toluene-metabolizing organisms and a generation time following the lag period of approximately 24 hours. In contrast, an inoculum of sucrose grown organisms into sucrose showed a lag period of approximately 1/2 hour and a generation time of 2 hours.

Qualitatively, the size of the cell pellets after centrifugation were approximately equal regardless of carbon source, and the protein assay revealed a protein content of 491.0ug/ml for sucrose grown organisms, 621.5ug/ml for

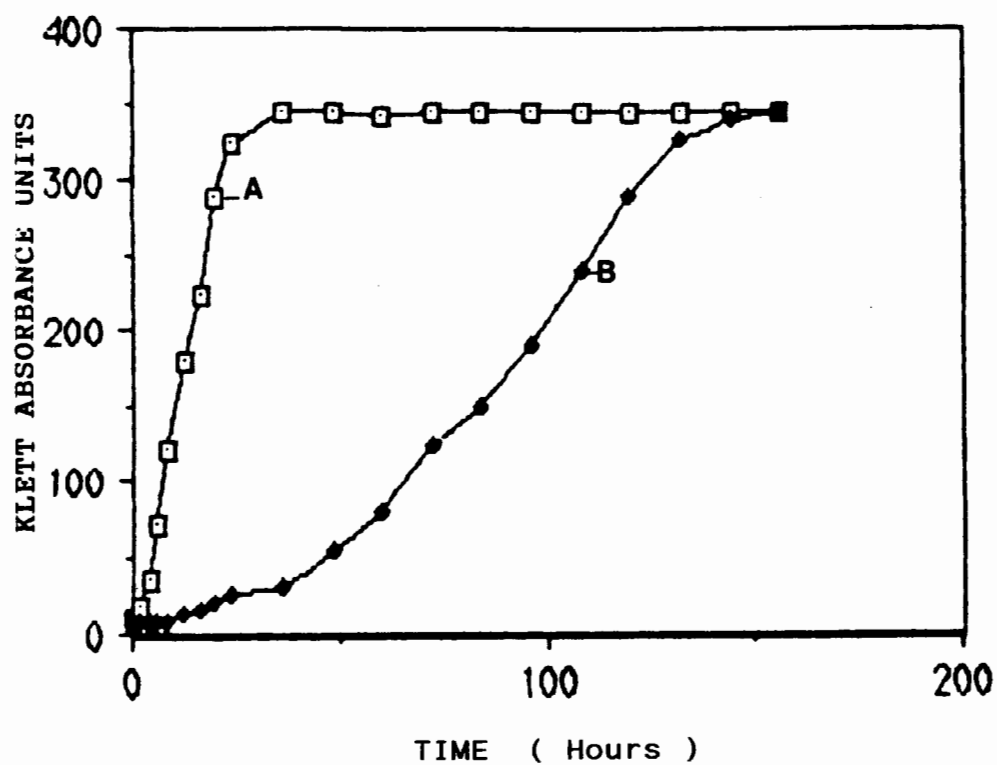


Figure 3. Growth of *P. putida* PC2P15. A) mBH with sucrose 0.1%; B) mBH with toluene vapor

sucrose and toluene grown bacteria, and 501.5ug/ml for sucrose and succinate grown organisms (Figure 4). This data represents the average of six identical experiments.

MORPHOLOGY

The morphology of PC2P15 grown with toluene as its sole source of carbon and energy was very different from the bacterium grown on mBH with sucrose as its sole carbon source. As shown in Figure 5, the toluene-grown organisms were approximately 1/3 the size of the cells grown in either sucrose or succinate. The toluene-grown cultures had a tendency to clump together around microdroplets of toluene. Individual bacteria could be observed "bending" around microdroplets of toluene and in direct contact with it. This observation occurred in both vapor-grown cells and in cultures to which toluene had been added directly. Nothing comparable to this clumping and bending phenomenon was observable in the sucrose-grown cells. The organisms were extremely motile in both toluene and sucrose cultures.

Electron microscopy (Figures 6,7,8) revealed the presence of membrane blebs in PC2P15 grown on toluene (Figure 8). High magnification (140,000x) revealed numerous blebs on the outer membrane of all cells examined. The blebs appeared to be surrounded by a bilayer membrane and, in some cases, to be "tethered" to the cell's outer membrane by this same bilayer membrane. Figure 6 shows this bilayer membrane.

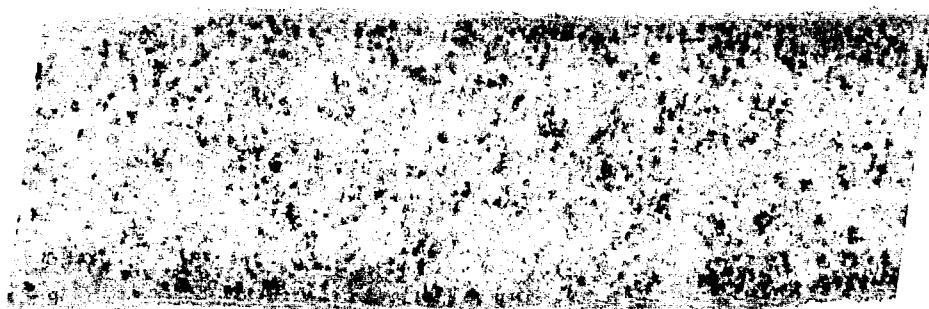
Figure 5. Light Micrographs of *P. putida* PC2P15. A) grown in toluene; B) grown in sucrose. Magnification: 1200X





Figure 6. Electron Micrograph of PC2P15 grown in toluene. Arrows indicate membrane blebs and extracellular vesicles. Magnification: 176,000X.

Figure 7. Electron Micrograph of PC2P15
grown in sucrose. Magnification: 176,000X.



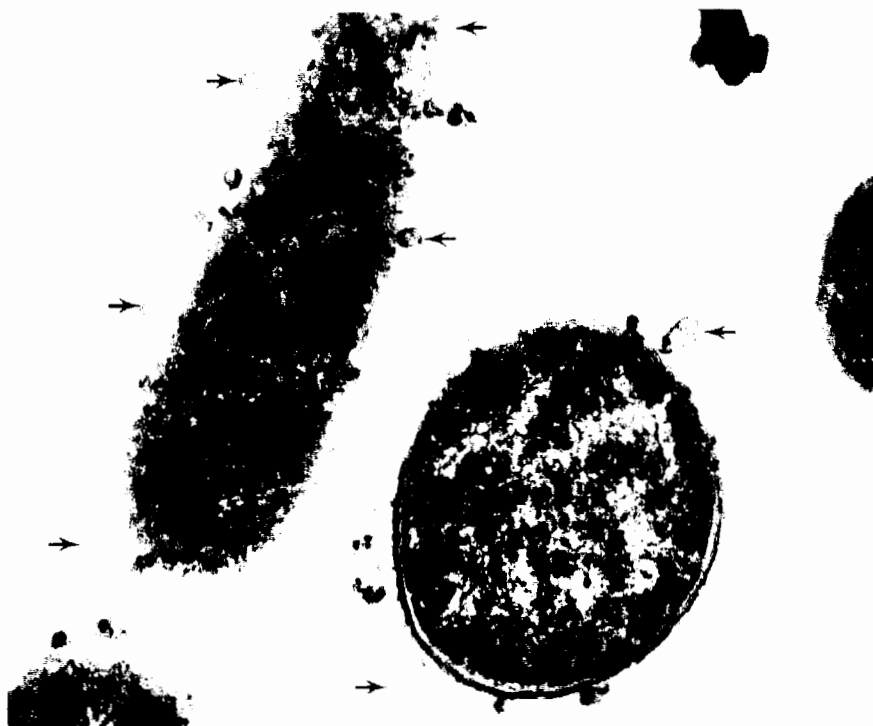


Figure 8. Electron Micrograph of PC2P15 grown in toluene. Arrows indicate membrane blebs and extracellular vesicles. Magnification: 140,000X.

The blebs were also been found intact and free-floating in the extracellular milieu. No such physical structures were present in the micrographs of the cells grown on sucrose (Figure 7).

Colony morphology on various agars was typical of *Pseudomonas putida*. The colonies were small, white, and opaque initially and, after 4 to 5 days of growth, appeared to have a nucleated center with an uneven, ragged edge. On TGYE and nutrient agar, the colonies became slimy after 3-4 days. The organisms were strongly positive for the production of siderophores on CAS agar (55).

Those colonies grown on agar on a Nucleopore filter as described in Materials and Methods did not exhibit this typical morphology. Although colonies were numerous and the bacteria grew readily, the colonies were very small and never nucleated. There was a lag period of approximately 48 hours when transferred to agar and exposed to toluene. Qualitatively, the colonies exposed to toluene appeared slimy. Microscopically, the toluene-grown cells were approximately 1/3 the size of cells grown on agar with sucrose as the carbon source. Colonies growing directly on the agar with sucrose as the carbon source and without the Nucleopore filter had typical *P.putida* morphology and did not become slimy. Colonies streaked onto a Nucleopore filter with no added carbon source showed no growth.

PC2P15 grew optimally at approximately 23⁰ C. Temperatures in excess of 32⁰ C showed increasingly slowed growth as a function of increased temperature and there was no growth at 42⁰ C as shown in Table II.

Aeration was important in the growth of PC2P15, especially when grown with toluene. At 100rpm, the growth rate was approximately 25% faster in those organisms growing on toluene when compared to organisms growing in toluene at 25rpm. Non-aerated organisms exhibited a significantly reduced growth rate when compared to those that were aerated. Aeration appeared to play no role in the growth rate until after the lag period had passed for the hydrocarbon-grown bacteria (ie: the lag period remained 48 hours).

Gross examination of the flask when the organisms were growing in toluene revealed the presence of a slime layer adhering to the sides of the flask. This slime layer was present in all actively growing, exponential phase cultures of toluene-grown organisms at the liquid/vapor interface. Living organisms could be easily recovered from this slime layer with a sterile loop. Centrifugation of liquid cultures grown with toluene were difficult to pellet and had a slimy appearance. Sucrose-grown controls never produced this slime layer and centrifugation produced a firm, non-slimy pellet.

When a cell pellet or drop of cell culture grown in toluene was placed on an oil surface it collapsed

immediately and floated at the surface. Cell pellets or drops of cultures of bacteria grown in sucrose did not collapse and sank within a few seconds. This suggested the production of an emulsifying agent or surfactant by PC2P15 growing in toluene.

FATTY ACID PROFILE

Fatty acid analysis of the toluene-grown organisms showed a number of significant changes in the membrane fatty acid profile. Figure 9 shows this data compared to the fatty acid profile for cells grown in nutrient broth. Several new fatty acids were synthesized by the toluene-grown bacteria, including octadecanoic and eicosanoic acids. Octadecanoic acid increased from 0% in the controls to 12.37% of the total fatty acid content in the toluene-grown cells. Conversely, there was a 36% decrease in undecanoic acid and a 56% decrease in trans-9 octadecenoic acid. There was a 100% decrease in nonadecanoic acid in the toluene-grown cells. Two major fatty acid components, dodecanoic and hexadecanoic acids, did not show significant changes when the toluene-grown cells were compared to the controls. The latter two fatty acids account for approximately 55% of the total fatty acid content of the cells.

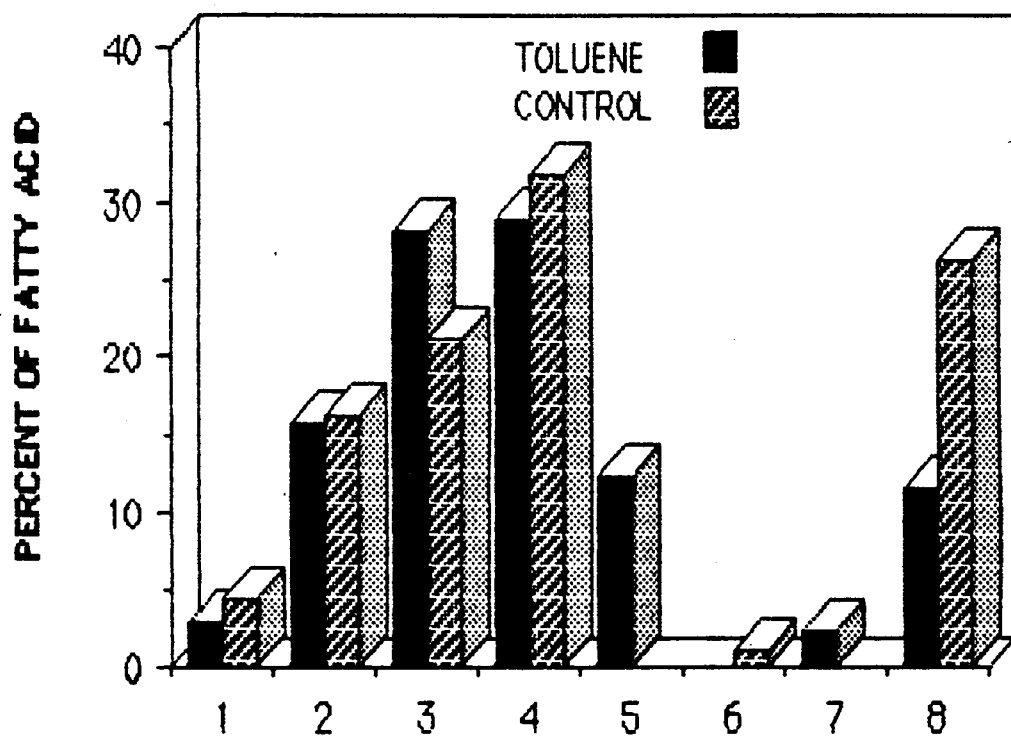


Figure 9. Fatty acid profile of PC2P15.
Columns: 1) undecanoic; 2) dodecanoic;
3) 9-hexadecenoic; 4) hexadecanoic;
5) octadecanoic; 6) nonadecanoic;
7) eicosanoic; 8) trans-9 octadecenoic.

METABOLIC AND PROTEIN ANALYSIS

SDS-PAGE analysis of lysed cells revealed a prominent 65kd band in toluene-grown cells and the appearance of new bands at 42kd and 44kd when compared to cells grown on sucrose, succinate or n-alkanes. Another group of low molecular weight bands (< 25kd) were present in toluene-grown cells, but not present in sucrose-, n-alkane-, or succinate-grown bacteria. A series of eight bands appeared in cells grown on n-alkanes that were not present in sucrose-, succinate-, or toluene-grown cells. A number of bands appeared in sucrose-grown bacteria that were not present in either toluene- or alkane-grown cells. Banding patterns for succinate-grown cells were not significantly different from those grown in toluene.

CATABOLITE REPRESSION

Studies were done to determine the rate of ^{14}C -sucrose uptake in cells in the presence of toluene, succinate, or toluene plus succinate. We also tested the effect of cyclic-AMP on ^{14}C -sucrose uptake with various combinations of these substrates.

Exponentially growing cells using toluene as the sole source of carbon and energy were inoculated in media with various carbon sources as shown in Table IV. The flasks were

TABLE IV
¹⁴C-SUCROSE UPTAKE IN PC2P15

FLASK #	FLASK CONTENTS**	CPM/mg protein*
1	SUCROSE	46,000
2	SUCROSE/TOLUENE	23,000
3	SUCROSE/TOLUENE/SUCCINATE	9,000
4	SUCROSE/TOLUENE/C-AMP	25,000
5	SUCROSE/C-AMP	44,000
6	SUCROSE/SUCCINATE	24,000
7	SUCROSE/SUCCINATE/C-AMP	23,000
8	SUCROSE/SUCCINATE/TOLUENE/C-AMP	10,000
9	FLASK 3 W/SUCROSE INOCULUM	12,000
10	FLASK 3 W/TOLUENE INOCULUM	10,000

* CPM ROUNDED TO NEAREST 1000.

**FINAL CONCENTRATIONS IN INDICATED FLASKS:

Sucrose 0.1%; Succinate 0.1%; C-AMP 1mM;

Toluene as vapor

incubated as described in Materials and Methods. The data in Figures 10 and 11 and the data in Table IV represent the average of 6 separate experiments.

When toluene or succinate was present in the medium containing sucrose (Flasks 2 and 6), the cpm/mg protein was decreased by approximately 50% compared to medium containing sucrose alone. Since succinate is reported to be a catabolite repressor in *Pseudomonas* (62), c-AMP was added to flasks containing sucrose and toluene (Flask 4), sucrose and succinate (Flask 7), sucrose alone (Flask 5) and sucrose, toluene, and succinate (Flask 8). It can be seen that c-AMP had no affect. It can also be seen that cultures grown with sucrose, succinate, and toluene without c-AMP (Flask 3) showed only 25% of the cpm/mg protein when compared to sucrose alone (Flask 1). Again, the addition of c-AMP (Flask 8) had no affect on uptake.

A separate experiment comparing an inoculum of sucrose-grown bacteria versus an inoculum of bacteria grown in toluene (Flasks 9 and 10) showed the same effect regardless of previous growth. The bacteria grown in sucrose had been transferred 4 times on sucrose prior to use as the inoculum.

Since the initial experiments were harvested at 144 hours, two identical experiments were done and the cells harvested at 24 and 48 hours. Time of harvest was shown to be of no importance and ^{14}C -sucrose was reduced by 50% and 75% as previously observed.

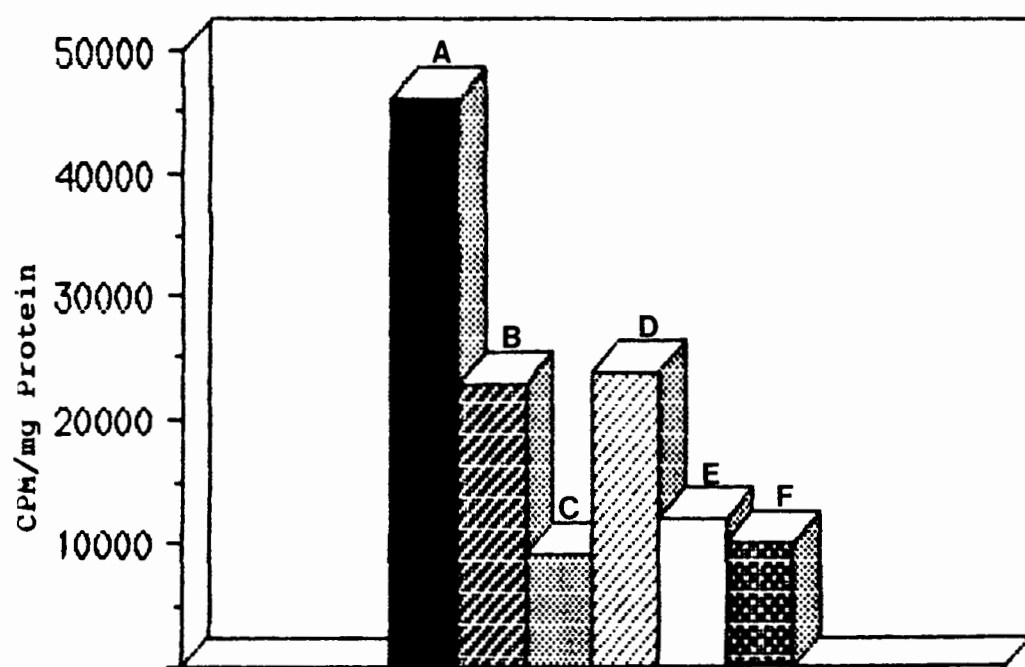


Figure 10. ^{14}C -sucrose uptake in *P. putida* PC2P15. A) Sucrose; B) Sucrose/Toluene; C) Sucrose/Toluene/Succinate; D) Sucrose/Succinate; E) Sucrose/Toluene/Succinate: sucrose inoculum; F) Sucrose/Toluene/Succinate: toluene inoculum

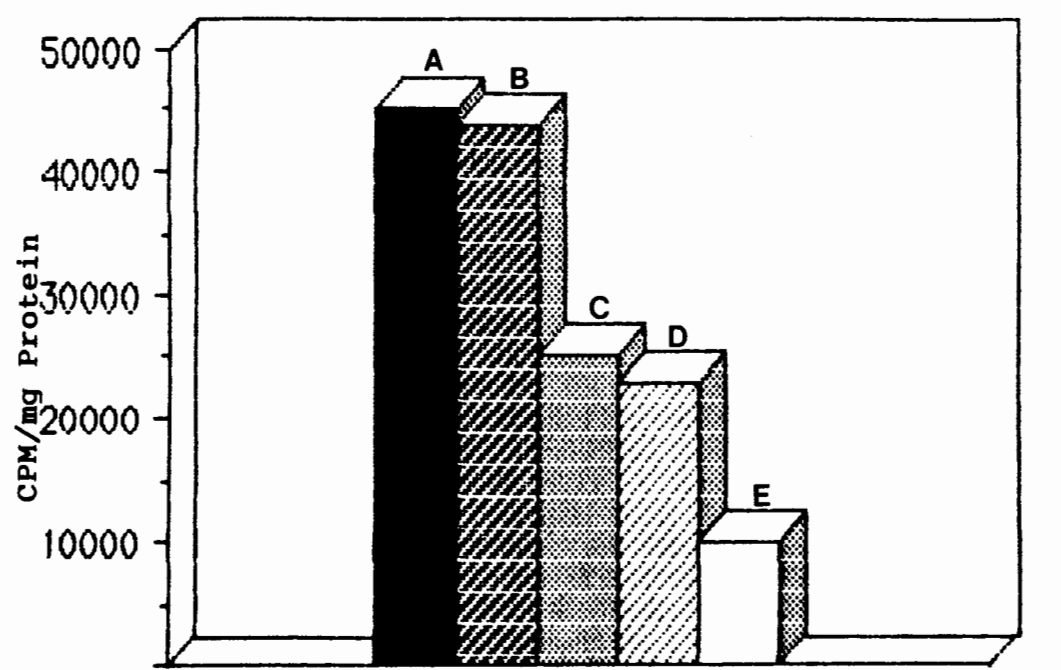


Figure 11. ^{14}C -Sucrose uptake in *P. putida* with cyclic-AMP added. A) Sucrose (no c-AMP); B) Sucrose; C) Sucrose/Toluene; D) Sucrose/Succinate; E) Sucrose/Succinate/Toluene.

Isotope balance showed that the amount of ^{14}C -sucrose remaining in the supernatant in toluene, succinate, or toluene/succinate cells was proportionally higher than the sucrose cells. In addition, the proportion of the original quantity of ^{14}C -sucrose not present in either the pellet or the supernatant was lower in toluene, succinate, and toluene/succinate cells when compared to sucrose controls. The ^{14}C not present in pellet or supernatant appears to have been respired by the cells in quantities that are approximately proportional to the observed uptake and utilization data in Table IV and in Figures 8 and 9. This suggests that ^{14}C -sucrose was not respired at the rate in sucrose/toluene, the sucrose/succinate, or the sucrose/succinate/toluene cells at the rate it was being respired in the sucrose controls.

DISCUSSION

MORPHOLOGY AND METABOLIC STRESS

The general morphology of organisms growing under laboratory conditions as opposed to those growing in natural environments has recently been reviewed by Roszak (30). In this excellent article, Roszak suggests that morphological changes, such as the reduction in size and slower growth rate that we have observed in our toluene grown organisms, may be closer to the natural state of bacteria in the environment. Roszak suggests that laboratory control organisms may, in fact, be abnormally large with an artificially high reproductive rate because of the unlimited access to nutrients and carbon sources rarely found in a natural setting.

Bacteria growing in a metabolically stressed environment may be viable but non-culturable (30). There are changes in fatty acid content, RNA, cellular DNA, and protein content in bacteria that are metabolically stressed or near starvation (31,32). It has been suggested that smaller size maximizes nutrient uptake by way of an increased surface area to volume ratio. In one study, DNA content in starved cells dropped to 5.2% of the original quantity after 98 days of starvation (31).

Bacteria in the natural environment may have a number of survival mechanisms, including the production of endospores. It has been hypothesized that bacteria unable to produce an endospore may have a non-spore survival stage in which they enter a period of dormancy with extremely low metabolic rates (30). In this state they are considered viable but non-culturable. Given an adequate nutrient supply and an easily assimilated substrate such as pyruvate, there is a rapid increase in intracellular ATP followed by an increase in motility, cellular DNA content, cellular size, and finally cell division following a lag period that is dependent on the type of organism present and the available substrate (30). This period of dormancy, in which the surviving bacteria (estimated at 2-4% of the original population) are believed to metabolize the cellular contents of lysed bacteria, has been termed cryptic growth (30).

The observations above could explain some of our findings with regard to the toluene oxidizing organism PC2P15. Organisms transferred from a minimal salts medium with sucrose to a minimal salts medium with toluene undergo a lag period of as long as 10 days before growth begins again. Microscopic examination of the first post-transfer exponential stage reveals that most of the organisms appear to have reduced their size by 1/2 to 2/3 compared to the size of the bacteria in the inoculum. This might indicate that PC2P15 does not use toluene as efficiently as sucrose

and reduces its size to increase its surface area to volume ratio. It may also indicate membrane damage as a result of toluene exposure and a resulting decrease in the efficiency of uptake of the toluene substrate.

In addition, we have experienced occasional difficulty regaining culturable organisms with the ability to metabolize toluene when sucrose-grown bacteria were transferred to toluene. This observation may be due in part to the fact that pseudomonads have a tendency to spontaneously lose plasmids over a period of time when grown in substrates other than those coded for on the plasmid. Summers (48) observed a failure to maintain plasmids in high copy numbers over time, and Cohen (49) detected a loss of plasmids in daughter cells following cell division. Keshavarz (56) noted a residual of only 1% of the population retained the m-toluate plasmid after 600 hours of growth in benzoate. We have found that increasing the initial inoculum size from 1ml/20ml to 10ml/20ml greatly enhances the recovery of toluene-oxidizing organisms from a sucrose culture. This suggests that only a small percentage of inoculated bacteria may have retained the ability to degrade toluene after a period of time using sucrose as a substrate. A second possibility is that there is an extended lag period when the alternate substrate is a hydrocarbon requiring a different degradative pathway. Lag times of up to 6 months have been reported for certain strains of bacteria (61). The

mechanism behind sudden exponential growth after a lag period of weeks or months remains in doubt.

A number of researchers have shown both interior and exterior morphological changes for organisms metabolizing hydrocarbons. In a species of *Acinetobacter*, Scott and others (50,51,52) found membrane-bound inclusion bodies in organisms metabolizing hexadecane. It is believed that these structural changes, which include the membrane-surrounded inclusion bodies, an intra-cytoplasmic membrane, and the production of extracellular vesicles, have a direct bearing on the utilization of hydrocarbons as sources of carbon and energy.

Electron micrographs of PC2P15 did not reveal the presence of either an intracytoplasmic membrane or hydrocarbon inclusion bodies. It did, however, reveal what appear to be blebs on the surfaces of cells grown in toluene as the sole source of carbon and energy. In addition to blebs on the surfaces of the cells, a number of these membrane-enclosed structures were observed free in the extracellular milieu. Kappeli (53) has noted similar extracellular vesicles in hexadecane-grown *Acinetobacter* and in a species of *Pseudomonas*. Because of the toxicity of toluene to most microorganisms, we initially suspected that this bleb phenomenon might have been a mechanism of membrane sloughing and repair because we observed an intact membrane beneath the blebs. We further speculated that the blebs

might have been the result of cell shrinkage. We now suggest that these blebs may be extracellular vesicles possibly involved in the emulsification and partitioning of toluene for use by the bacterium.

Experiments currently in progress in our laboratory in which sterile filtered medium from an exponentially growing culture of PC2P15 is used as the starting media for a fresh inoculum of PC2P15, have shown promising preliminary results. The lag time for toluene-grown organisms is reduced from approximately 48 hours to approximately 6 hours. This may indicate the presence in the medium of a bacterial product or products that assists with the partitioning, emulsification, or uptake of the toluene substrate. If this is the case, it would help to explain the characteristic 48 hour lag phase in transfers of exponentially-grown toluene organisms into toluene. Although the toluene-grown organisms have a much slower growth rate than the sucrose-grown cells, this does not adequately explain the 48 hour period of no growth or minimal growth we have observed. We suggest that these organisms may "condition" the medium by creating a microemulsion of toluene through the release of an as yet unidentified agent or agents into the medium by way of the observed membrane vesicles. Other pseudomonads are known to release a rhamnolipid emulsifying agent when grown on alkanes to increase the solubility of the alkane (53). There is reason to suggest that a similar mechanism to

solubilize toluene may operate in PC2P15. The slimy biofilm observed at the liquid/vapor interface where those organisms were growing in toluene may have been part of the visible manifestation of those emulsifying agents. The biofilm may also have served to allow the organisms to adhere to the sides of the flasks at the liquid/vapor interface where they were continuously bathed in nutrients from the agitation of the rotating shaker. We are now attempting to identify both the nature of the membrane blebs and of the biofilm produced by the bacteria.

CYTOPLASMIC PROTEINS

Total protein in the cell pellet of organisms grown on toluene was approximately the same as the protein content in the cell pellet of organisms grown on sucrose, sucrose and toluene, or sucrose, toluene, and succinate. These results support our observation and the observation of other researchers (31) that, although the toluene-grown cells were smaller than sucrose grown cells, total protein content was approximately equal in the cell pellets and that the number of cells in the population was equal to or higher than the sucrose grown cells. It was important to ascertain cell density by both the turbidity of the cultures and by assay of total protein content to properly interpret our findings in the sucrose uptake and SDS-PAGE studies.

SDS-PAGE of the cell lysate revealed that sucrose-grown cells produced a banding pattern significantly different from cells grown on toluene. Prominent bands of about 65kd, 42kd, 44kd, and several bands below 25kd that were not present in either the sucrose or the alkane grown cells were observed in cells grown on the aromatics toluene, benzene, and phenol. These findings suggest the activation of genes involved in the acquisition or degradation of aromatics. It is reasonable to assume that these and other new bands present in the aromatic grown cells were enzymes or structural proteins involved in either the degradation of toluene, benzene and phenol or as a result of exposure to these substances. In the case of phenol, we might expect to find a monooxygenase being used due to the presence of the hydroxyl group present on the ring, but a banding difference between toluene and phenol was not seen. However, the use of the dioxygenase system, resulting in a triple hydroxylation followed by a dehydration to form the catechol, cannot be ruled out at present.

A series of 8 bands appeared in the alkane-grown cells that were not present in the aromatic- or sucrose-grown cells. When the cells were grown with sucrose and toluene, the banding pattern was a composite of the banding patterns of the two separate carbon sources but the sucrose bands were less intense. Lysates of organisms grown in toluene, sucrose, and succinate showed no change in banding from that

of the sucrose/toluene grown cells. This result might be expected from the addition of succinate, a Krebs cycle intermediate, that would not necessarily require the induction of new catabolic enzymes.

METABOLISM OF MULTIPLE CARBON SOURCES

Our data suggest that both sucrose and toluene are catabolized by inducible enzyme systems, and that PC2P15 has the capacity to use both carbon sources at the same time. A recent study (54) indicated that *Pseudomonas* strain JS6 has the capacity to simultaneously metabolize both chlorobenzene and toluene which requires the concurrent induction of two metabolic pathways. The former entails an ortho-fission pathway and the latter uses a meta-fission pathway. It was previously believed that these two pathways were incompatible. Another organism, *Alcaligenes eutrophus* JMP 134, has been shown to use three pathways at once (54). This organism utilizes an ortho-, modified ortho-, and a meta-fission pathway simultaneously .

Our ^{14}C -sucrose uptake studies have indicated that PC2P15 may utilize toluene, sucrose, and succinate simultaneously. It is known that the membrane protein, succinic dehydrogenase, is produced constitutively in many bacteria, including the pseudomonads. The metabolic pathway for sucrose, if not constitutive, would require the induction of enzymes like phosphotransferase and others

necessary for sucrose uptake and utilization. It is not known whether the enzymes of the Entner-Doudoroff pathway are produced constitutively in this particular strain after a period of growth on a toluene substrate requiring a different degradative pathway.

Succinate, which is known to be a catabolite repressor in *P. putida* (62), depressed sucrose uptake approximately 50% in cells exposed to both sucrose and succinate. It is unclear why the presence of cyclic-AMP did not derepress sucrose uptake in cultures of sucrose and succinate or in cultures of sucrose and toluene if either succinate or toluene were acting as catabolite repressors.

Our results indicated an activation of both the toluene and sucrose enzyme systems in organisms with both toluene and sucrose as simultaneously available carbon sources. When succinate was added to toluene and sucrose cultures, ^{14}C -sucrose uptake was depressed by an additional 20-25% of that with sucrose and toluene alone. This suggests the simultaneous use of three distinct carbon sources in the same organism. It is possible that a feedback mechanism operates in PC2P15 to decrease the uptake of ^{14}C -sucrose in the presence of abundant Krebs intermediates derived from the uptake and utilization of toluene and/or succinate. Our data suggest that the enzymes of the Entner-Doudoroff pathway, the proteins required for succinate uptake and

utilization, and the pathway necessary for the catabolism of toluene are simultaneously operative in PC2P15.

Another possibility is that, in any given population of bacteria, individual bacteria may be utilizing only one of the two or three available carbon sources. For example, in a flask of PC2P15 exposed to toluene, succinate, and sucrose, 25% of the bacteria may be using sucrose, 50% may be using toluene, and 25% using succinate. This possibility cannot be dismissed although our consistent results suggest the simultaneous use of multiple carbon sources by PC2P15.

Studies are planned in our laboratory that we hope will further elucidate how PC2P15 utilizes multiple carbon sources and the mechanisms involved. In addition, we hope to use monoclonal antibodies to the mono- and dioxygenases and to cis-toluene dihydrodiol dehydrogenase to elucidate both the roles and locations of these enzymes intracellularly or in the membrane vesicles.

CONCLUSIONS AND SUMMARY

A strain of *Pseudomonas* has been isolated from a bioremediation pond and identified as a *Pseudomonas putida* which we have designated strain PC2P15. This organism has exhibited the ability to use a number of aliphatic hydrocarbons, aromatics, and other substrates as the sole sources of carbon and energy.

Our results have shown significant changes in the fatty acid content, the synthesis of proteins, the growth rate, and the morphology of these bacteria when using toluene as the sole source of carbon and energy when compared to cells grown in sucrose. We have observed membrane blebs or extracellular vesicles attached to the outer surface of cells and also in the extracellular milieu of toluene-grown organisms. We believe these blebs may be involved in the emulsification and partitioning of the toluene substrate for use by the bacteria.

Our results also indicated the simultaneous use of multiple carbon sources by PC2P15. ^{14}C -sucrose uptake was decreased by 50% when cells were grown with sucrose and toluene or in sucrose with succinate. Our results have also shown an additional 20-25% drop in sucrose uptake when succinate was added to a sucrose/toluene culture.

In light of the increasing toxic contamination of both soil and water in the world today, science is striving to understand the role of microorganisms in the degradation of xenobiotics. Even though many bacteria possess the metabolic capacity to use hydrocarbons, the pseudomonads have a particularly wide range of metabolic abilities which are of interest to microbiologists.

Although a great deal of information has been gathered over the past several decades on the physiology, biochemistry, and genetics of biodegradative bacteria, much more work still lies ahead. It is not enough to study these organisms in the laboratory. To fully understand how these cells operate, they must be studied *in situ*. Because isolated organisms in the laboratory probably function in a very different manner from those in the environment, a complete understanding of the bioremediation of contaminated soils and water by microorganisms will come only from the study of these organisms in their natural habitats and with proper consideration given to their relationships to other organisms and to the abiotic conditions present in their environment. A complete and long term effort is necessary to study these organisms in both amended and unamended environments *in situ* to compliment the information obtained from the laboratory.

It is of interest that our isolate may have the capability to metabolize multiple carbon sources simul-

taneously. In the environment, concurrent metabolism of several carbon sources might have a survival advantage for those organisms that have evolved these abilities. If our isolate also has the capability to secrete emulsifying agents or enzymes into the environment in a manner similar to the release of iron-chelating siderophores, it would gain a definite evolutionary advantage over those bacteria without these heritable traits.

Because an entire bacterial community can exist and evolve on a single grain of soil to which a particular substrate has adhered, it is difficult to follow the progress of a single population of organisms *in situ*. We hope that, over the next few years, science will develop the technical expertise to observe and understand the actions and interactions of microorganisms *in situ* and gain insights into their exceedingly complex and vital role in the degradation of naturally occurring substances and xenobiotics in the environment.

REFERENCES

- (1) Eastern Research Group, Inc. (1989). Alaskan Oil Spill Bioremediation Project. In: *Ecol.Res.Ser. U.S. Environmental Protection Agency*.
- (2) Hill, D.L., et al (1989). *Biochem.Biotechnol.* 20-21, 233-244.
- (3) von Wedel, R.J., et al (1988). *Water Science Technol.* 20:11-12, 501-503.
- (4) Ridgway, H.F., et al (1990). *Applied Microbiology Biotechnology.* 34:2, 248-252.
- (5) Leahy, J.G. and R.R. Colwell (1990). *Microbiol. Rev.* 54:3, 305-315.
- (6) Premuzik, E.T. and M.S. Lin (1991). *Resources Conservation and Recycling.* 5:2-3, 167-182.
- (7) VanDenmark, Paul J. and B.J. Batzung (1987). *The Microbes.* Benjamin Cummings Pub. Co.. 497-499.
- (8) Blumer, M. and W.W. Youngblood (1975) *Science.* vol. 188. 53-55.
- (9) Hites, R.A., et al (1977). *Science.* vol. 198. 829-831.
- (10) Reineke, W. (1984). In: *Microbial Degradation of Organic Compounds.* D.T. Gibson (ed.). 319-360.
- (11) Chapman, P.J. (1988). In: *Environmental Biotechnology: Reducing the Risks from Environmental Pollution Through Biotechnology.* G.S. Omenn (ed.). 81-95.
- (12) Timmis, K.N., et al (1988). In: *Environmental Biotechnology: Reducing the Risks from Environmental Pollution Through Biotechnology.* G.S. Omenn (ed.). 61-79.
- (13) Ihoue, Akira and Koki Horikoshi (1989). *Nature.* 338, 264- 266.
- (14) Dagley, Stanley (1978). *Q. Rev. Biophys.* 11, 577-602.

- (15) Anderson, J.J. and S. Dagley (1981). *J. Bacteriol.* 146, 291-297.
- (16) Gibson, D.T., et al (1968). *Biochemistry.* 7:2653.
- (17) Gibson, D.T., et al (1970). *Biochemistry.* 9:1631.
- (18) Hogn, T., et al (1972). *Eur. J. Biochem.* 30:369.
- (19) Dagley, Stanley (1986). In: *The Bacteria: A Treatise on Structure and Function.* J.R. Sokatch (ed.). Academic Press 527-555.
- (20) Stanier, R.Y., et al (1966). *J. Gen. Microbiol.* 43, 159-271.
- (21) Slater, J.H. and D.J. Hardman (1982). In: *Experimental Microbial Ecology.* R.G. Burns and J.H. Slater (eds.). Blackwell, Oxford. 255-274.
- (22) Sato, T., et al (1963). *J. Biochem.* 52:23.
- (23) Jerina, D.M., et al (1968). *Arch. Biochem. Biophys.* 128:176.
- (24) Dagley, Stanley (1986). In: *The Bacteria: A Treatise on Structure and Function.* J.R. Sokatch (ed.). Academic Press. 527-555.
- (25) *Methods in Enzymology.* (1974). S. Fleischer and L. Packer (eds.). Vol. 31. Academic Press.
- (26) *Methods in Enzymology.* (1984). W.B. Jacoby (ed.). Vol. 104. Academic Press.
- (27) Pakrasi, T., et al (1979). *Proc.Nat.Acad.Sci.* Vol. 82. 6903-6907.
- (28) Smith P.K., et al (1985). *Anal. Biochem.* 150: 76-85.
- (29) Jain, D.K., et al (1991). *Journal of Microbiological Methods.* 13:4, 271-280.
- (30) Roszak, D.B. and R.R. Colwell (1987). *Microbiological Reviews.*
- (31) Moyer, Craig L. and Richard I. Morita (1989). *Applied and Environmental Microbiology.* 55:10, 2710-2716.
- (32) Linder, Katherine and James D. Oliver (1989). *Applied and Environmental Microbiology.* 55:11, 2837-2842.

- (33) Yen, Kwang-Nru, et al (1991). *J. Bacteriol.* 173:17, 5315-5327.
- (34) Rogers, J.E. and D.T. Gibson (1977). *J. Bacteriol.* 130:3, 1117-1124.
- (35) Collingsworth, William L., et al (1973). *J. Bacteriol.* 113:2, 922-931.
- (36) Whited, Gregory M. and David T. Gibson (1991). *J. Bacteriol.* 173:9, 3010-3016.
- (37) Crutcher, Stephen E. and Phillip J. Geary (1979). *Biochemical Journal.* 177, 393-400.
- (38) Subramanian, V., et al (1981). *The Journal of Biological Chemistry.* 256:6, 2723-2730.
- (39) Gibson, D.T., et al (1970). *Biochemistry.* 9:7, 1631-1635.
- (40) Gibson, D.T., et al (1970). *Biochemistry.* 9:7, 1626-1630.
- (41) Madsen, Eugene L., et al (1991). *Science.* 252:10, 830-833.
- (42) King, J.M.H., et al (1990). *Science.* 251:17, 778-781.
- (43) Claxton, Larry D., et al (1991). *Nature.* 353, 24-25.
- (44) Brumsbach, F.R. and W. Reineke (1990). *Applied Microbiology and Biotechnology.* 34:2, 248-252.
- (45) Heitkamp, M.A., et al (1990). *Applied and Environmental Microbiology.* 56:10, 2967-2973.
- (46) Pettigrew, C.A., et al (1990). *Applied and Environmental Microbiology.* 57:7, 2036-2045.
- (47) Ramos, Gonzalez M.I., et al (1991). *Applied and Environmental Microbiology.* 57:10, 3020-3027.
- (48) Summers, D.K. and D. Sherratt (1984). *Cell.* 36, 1097-1103.
- (49) Cohen, S.N., et al (1985). In *Plasmids in Bacteria.* D.R. Helinski, S.N. Cohen, D.B. Clewell, D.A. Jackson, and A. Hollaender (eds.). Plenum. 189-214.
- (50) Scott, Christina C.L. and W.R. Finnerty (1976). *J. Bacteriol.* 127:1, 481-489.

- (51) Scott, Christina C.L., et al (1976). *J. Bacteriol.* 127:1, 469-480.
- (52) Scott, Christina C.L. and W.R. Finnerty (1976). *Journal of General Microbiology.* 94, 342-350.
- (53) Kappeli, O. and W.R. Finnerty (1979). *J. Bacteriol.* 140:2, 707-712.
- (54) Pettigrew, Charles A., et al (1991). *Applied and Environmental Microbiology.* 57:1, 157-162.
- (55) Schwyn, Bernhard and J.B. Neilands (1987). *Analytical Biochemistry.* 160, 47-56.
- (56) Keshavarz, T., et al (1985). *Journal of General Microbiology.* 131, 1193-1203.
- (57) Goldstein, Rebecca M., et al (1985). *Applied and Environmental Microbiology.* 50:4, 977-983.
- (58) Alvarez, Pedro J.J. and Timothy M. Vogel (1991). *Applied and Environmental Microbiology.* 57:10, 2981-2985.
- (59) Sander, Peter, et al (1991). *Applied and Environmental Microbiology.* 57:6, 1430-1440.
- (60) Lindstrom, Jon E., et al (1991). *Applied and Environmental Microbiology.* 57:9, 2514-2522.
- (61) Linkfield, Timothy G., et al (1989). *Applied and Environmental Microbiology.* 55:11, 2773-2778.
- (62) *The Microbial World* 4th Ed. (1976). R.Y. Stanier, E.A. Adelberg, and J.L. Ingram (eds.). Prentice-Hall. pp 257.
- (63) *Oil in the Sea: Inputs, Fates, and Effects* (1985). National Academy Press. Washington, D.C.. p 47.